

PCT09

## RAW SEQUENCE LISTING DATE: 04/21/2003 PATENT APPLICATION: US/09/913,770A TIME: 13:43:13

Input Set : N:\AMC\SEQ068rev.txt

Output Set: N:\CRF4\04212003\I913770A.raw

```
3 <110> APPLICANT: Takeda Chemical Industries, Ltd.
      5 <120> TITLE OF INVENTION: Novel G Protein Coupled Receptor Protein and Its DNA
      7 <130> FILE REFERENCE: 2590WOOP
C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/913,770A
C--> 9 <141> CURRENT FILING DATE: 2001-08-17
     9 <150> PRIOR APPLICATION NUMBER: JP 11-041336
     10 <151> PRIOR FILING DATE: 1999-02-19
     12 <150> PRIOR APPLICATION NUMBER: JP 11-125768
    13 <151> PRIOR FILING DATE: 1999-05-06
                                                                  Does Not Comply
    15 <160> NUMBER OF SEQ ID NOS: 4
                                                              Corrected Diskette Needed
    17 <210> SEO ID NO: 1
                                                                pr 2-3
    18 <211> LENGTH: 340
    19 <212> TYPE: PRT
    20 <213> ORGANISM: Human
    22 <400> SEOUENCE: 1
    23 Met Asn Pro Phe His Ala Ser Cys Trp Asn Thr Ser Ala Glu Leu Leu
    25 Asn Lys Ser Trp Asn Lys Glu Phe Ala Tyr Gln Thr Ala Ser Val Val
                    20
                                         25
    27 Asp Thr Val Ile Leu Pro Ser Met Ile Gly Ile Ile Cys Ser Thr Gly
                35
                                     40
    29 Leu Val Gly Asn Ile Leu Ile Val Phe Thr Ile Ile Arg Ser Arg Lys
    31 Lys Thr Val Pro Asp Ile Tyr Ile Cys Asn Leu Ala Val Ala Asp Leu
    33 Val His Ile Val Gly Met Pro Phe Leu Ile His Gln Trp Ala Arg Gly
                        85
                                             90
    35 Gly Glu Trp Val Phe Gly Gly Pro Leu Cys Thr Ile Ile Thr Ser Leu
                                        105
    37 Asp Thr Cys Asn Gln Phe Ala Cys Ser Ala Ile Met Thr Val Met Ser
               115
                                    120
                                                        125
    39 Val Asp Arg Tyr Phe Ala Leu Val Gln Pro Phe Arg Leu Thr Arg Trp
                               135
                                                    140
    41 Arg Thr Arg Tyr Lys Thr Ile Arg Ile Asn Leu Gly Leu Trp Ala Ala
                           150
                                                155
    43 Ser Phe Ile Leu Ala Leu Pro Val Trp Val Tyr Ser Lys Val Ile Lys
                                            170
    45 Phe Lys Asp Gly Val Glu Ser Cys Ala Phe Asp Leu Thr Ser Pro Asp
                   180
                                        185
    47 Asp Val Leu Trp Tyr Thr Leu Tyr Leu Thr Ile Thr Thr Phe Phe
                                   200
                                                        205
    49 Pro Leu Pro Leu Ile Leu Val Cys Tyr Ile Leu Ile Leu Cys Tyr Thr
```

215

210

50

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```
51 Trp Glu Met Tyr Gln Gln Asn Lys Asp Ala Arg Cys Cys Asn Pro Ser
     52 225
                            230
                                                 235
     53 Val Pro Lys Gln Arg Val Met Lys Leu Thr Lys Met Val Leu Val Leu
                        245
                                             250
     55 Val Val Val Phe Ile Leu Ser Ala Ala Pro Tyr His Val Ile Gln Leu
                    260
                                         265
     57 Val Asn Leu Gln Met Glu Gln Pro Thr Leu Ala Phe Tyr Val Gly Tyr
     58
                                     280
                                                         285
     59 Tyr Leu Ser Ile Cys Leu Ser Tyr Ala Ser Ser Ser Ile Asn Pro Phe
                                295
                                                     300
     61 Leu Tyr Ile Leu Leu Ser Gly Asn Phe Gln Lys Arg Leu Pro Gln Ile
                            310
                                                 315
     63 Gln Arg Arg Ala Thr Glu Lys Glu Ile Asn Asn Met Gly Asn Thr Leu
                        325
                                             330
     65 Lys Ser His Phe
     66
                    340
     68 <210> SEQ ID NO: 2
     69 <211> LENGTH: 1023
     70 <212> TYPE: DNA
     71 <213> ORGANISM: Human
     73 <400> SEQUENCE: 2
     74 atgaatccat ttcatgcatc ttgttggaac acctctgccg aacttttaaa caaatcctgg
                                                                               60
     75 aataaagagt ttgcttatca aactgccagt gtggtagata cagtcatcct cccttccatg
                                                                              120
     76 attgggatta tctgttcaac agggctggtt ggcaacatcc tcattgtatt cactataata
                                                                              180
     77 agatccagga aaaaaacagt ccctgacatc tatatctgca acctggctgt ggctgatttg
                                                                              240
     78 gtccacatag ttggaatgcc ttttcttatt caccaatggg cccgaggggg agagtgggtg
                                                                              300
     79 tttggggggc ctctctgcac catcatcaca tccctggata cttgtaacca atttgcctgt
                                                                              360
     80 agtgccatca tgactgtaat gagtgtggac aggtactttg ccctcgtcca accatttcga
                                                                              420
     81 ctgacacgtt ggagaacaag gtacaagacc atccggatca atttgggcct ttgggcagct
                                                                              480
     82 teetttatee tggeattgee tgtetgggte taetegaagg teateaaatt taaagaeggt
                                                                              540
     83 gttgagagtt gtgcttttga tttgacatcc cctgacgatg tactctggta tacactttat
                                                                              600
     84 ttgacgataa caacttttt tttccctcta cccttgattt tggtgtgcta tattttaatt
                                                                              660
     85 ttatgctata cttgggagat gtatcaacag aataaggatg ccagatgctg caatcccagt
                                                                              720
     86 qtaccaaaac agagagtgat gaagttgaca aagatggtgc tggtgctggt ggtagtcttt
                                                                              780
     87 atcctgagtg ctgcccctta tcatgtgata caactggtga acttacagat ggaacagccc
                                                                              840
     88 acactggcct tctatgtggg ttattacctc tccatctgtc tcagctatgc cagcagcagc
                                                                              900
     89 attaaccett ttetetacat eetgetgagt ggaaatttee agaaaegtet geetcaaate
                                                                              960
     90 caaagaagag cgactgagaa ggaaatcaac aatatgggaa acactctgaa atcacacttt
                                                                             1020
     91 tag
                                                                             1023
     93 <210> SEQ ID NO: 3
     94 <211> LENGTH: 24
     96 <213 ORGANISM Artificial Sequence Ill Item // on Euro Summary Sheet
98 <220> FEATURE:
     99(<223) OTHER INFORMATION:
W--> 101 <400> 3
     102 atgaatccat ttcatgcatc ttgt
                                                             24
     104 <210> SEQ ID NO: 4
```

8.3

105 <211> LENGTH: 25

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/913,770A

DATE: 04/21/2003 TIME: 13:43:13

Input Set : N:\AMC\SEQ068rev.txt

Output Set: N:\CRF4\04212003\I913770A.raw see tem 1/on Enon Summony Sheet

106 <212> TYPE: DNA

107 <213 ORGANISM: Artificial Sequence 109 <220> FEATURE:

110 <223 > OTHER INFORMATION:

W--> 112 <400> 4

113 ctaaaagtgt gatttcagag tgttt

25

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/913,770A

DATE: 04/21/2003 TIME: 13:43:14

1 11ME: 13:

Input Set : N:\AMC\SEQ068rev.txt

Output Set: N:\CRF4\04212003\I913770A.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No

L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:101 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:99 L:112 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:4,Line#:110 STATISTICS SUMMARY

PATENT APPLICATION: US/09/913,770A T

DATE: 04/21/2003 TIME: 13:43:14

Input Set : N:\AMC\SEQ068rev.txt

Output Set: N:\CRF4\04212003\I913770A.raw

Application Serial Number: US/09/913,770A

Alpha or Numeric or Xml: Numeric

Application Class:

Application File Date: 08-17-2001

Art Unit: PCT09

Software Application:

Total Number of Sequences: 4

Total Nucleotides: 1072 Total Amino Acids: 340 Number of Errors: 0 Number of Warnings: 2 Number of Corrections: 2

## MESSAGE SUMMARY

258 W: 2 (Mandatory Feature missing)

270 C: 1 (Current Application Number differs)

271 C: 1 (Current Filing Date differs)